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Reviewer: Keisha Douglas

Timestamp: [year=2008; month=4; day=15; hr=17; min=17; sec=21; ms=469;]

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Application No: 10536802 Version No: 2.1

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Finished: 2008-04-15 17:15:24.992
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Total Errors: 0
No. of SeqIDs Defined: 10
Actual SeqID Count: 10

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SEQUENCE LISTING

<110> Japan Science and Technology Agency
Tsukasa SEYA
Misako MATSUMOTO
Hiroyuki OSHIUMI

<120> Novel Adaptor Protein that Binds to Mammalian Toll-Like Receptor 3,
and Gene Thereof

<130> 1035-591 / A211-02/PCT

<140> 10/536,802

<141> 2005-09-22

<150> PCT/JP2003/014854

<151> 2003-11-20

<150> JP 2002-349015

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Thr Pro Arg Pro Gly Cys Gln Gly Gln Asp Leu Leu His Ala Met Val
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Pro Glu Pro Val Pro Gly Gly Cys Gln Glu Pro Glu Glu Met Ser Trp	
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Leu Ser Val Glu Asp Thr Thr Ser Pro Asn Thr Lys Pro Cys Pro Pro	
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Gln Val Pro Gly Arg Gly Glu Leu Ser Cys Leu Gln Asp Ala Ile Asp	
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His Ser Ala Phe Ile Ile Leu Leu Leu Thr Ser Asn Phe Asp Cys Arg	
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Pro Ala Gln Leu Ser Ser Asp Thr Ala Ser Leu Leu Ser Gly Leu Val	
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cgg ctg gac gaa cac tcc cag atc ttc gcc agg aag gtg gcc aac acc	1643
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Ser Tyr Leu Ser Tyr Gln Ala Gln Met Glu Gln Leu Gln Val Ala Phe			
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Pro Phe Gly Gly Gln Val Pro Leu Gly Ala Pro Pro Pro Phe Pro Thr			
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Trp Pro Gly Cys Pro Gln Pro Pro Pro Leu His Ala Trp Gln Ala Gly			
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Thr Pro Pro Pro Pro Ser Pro Gln Pro Ala Ala Phe Pro Gln Ser Leu			
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ccc ttc ccg cag tcc cca gcc ttc cct acg gcc tca ccc gca ccc cct			2075
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Gln Ser Pro Gly Leu Gln Pro Leu Ile Ile His His Ala Gln Met Val			
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Gln Leu Gly Leu Asn Asn His Met Trp Asn Gln Arg Gly Ser Gln Ala			
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Pro Glu Asp Lys Thr Gln Glu Ala Glu			
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85 90 95

Ala Arg Leu Tyr His Leu Leu Ala Glu Glu Lys Leu Cys Pro Ala Ser
100 105 110

Leu Arg Asp Val Ala Tyr Gln Glu Ala Val Arg Thr Leu Ser Ser Arg
115 120 125

Asp Asp His Arg Leu Gly Glu Leu Gln Asp Glu Ala Arg Asn Arg Cys
130 135 140

Gly Trp Asp Ile Ala Gly Asp Pro Gly Ser Ile Arg Thr Leu Gln Ser
145 150 155 160

Asn Leu Gly Cys Leu Pro Pro Ser Ser Ala Leu Pro Ser Gly Thr Arg
165 170 175

Ser Leu Pro Arg Pro Ile Asp Gly Val Ser Asp Trp Ser Gln Gly Cys
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Ser Leu Arg Ser Thr Gly Ser Pro Ala Ser Leu Ala Ser Asn Leu Glu
195 200 205

Ile Ser Gln Ser Pro Thr Met Pro Phe Leu Ser Leu His Arg Ser Pro
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225 230 235 240

Glu Pro Val Pro Gly Gly Cys Gln Glu Pro Glu Glu Met Ser Trp Pro
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Pro Val Lys Asn Pro Cys Ser Val Lys Asp Gln Thr Pro Leu Gln Leu	325	330	335
Ser Val Glu Asp Thr Thr Ser Pro Asn Thr Lys Pro Cys Pro Pro Thr	340	345	350
Pro Thr Thr Pro Glu Thr Ser Pro Pro Pro Pro Pro Pro Pro Ser	355	360	365
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Ser Leu Glu Ser Ser Ser Glu Gln Lys Phe Tyr Asn Phe Val Ile Leu	385	390	395 400
His Ala Arg Ala Asp Glu His Ile Ala Leu Arg Val Arg Glu Lys Leu	405	410	415
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Val Pro Gly Arg Gly Glu Leu Ser Cys Leu Gln Asp Ala Ile Asp His	435	440	445
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Gly Ser Pro Asp Cys Val Ile Pro Phe Leu Pro Leu Glu Ser Ser Pro	485	490	495
Ala Gln Leu Ser Ser Asp Thr Ala Ser Leu Leu Ser Gly Leu Val Arg	500	505	510
Leu Asp Glu His Ser Gln Ile Phe Ala Arg Lys Val Ala Asn Thr Phe	515	520	525
Lys Pro His Arg Leu Gln Ala Arg Lys Ala Met Trp Arg Lys Glu Gln	530	535	540
Asp Thr Arg Ala Leu Arg Glu Gln Ser Gln His Leu Asp Gly Glu Arg	545	550	555 560
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Ser Pro Gly Leu Gln Pro Leu Ile Ile His His Ala Gln Met Val Gln
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